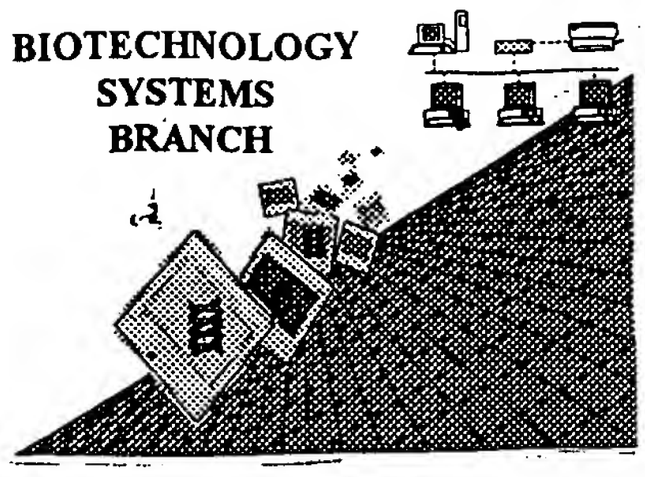
S. Foley

RAW SEQUENCE LISTING ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/509,712A

Source: 1600 RUSH

Date Processed by STIC: 11/7/2001

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

1) **INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,**
2) **TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.
PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)
PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE **CHECKER VERSION 3.0 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.
Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

http://www.uspto.gov/web/offices/pac/checker

1600

RAW SEQUENCE LISTING DATE: 11/07/2001
PATENT APPLICATION: US/09/509,712A TIME: 13:41:54

Input Set : N:\Crf3\09192001\I509712A.raw
Output Set: N:\CRF3\11072001\I509712A.raw

see pp 1-5

```
1 <110> APPLICANT: Rubin, Donald H.
2       Organ, Edward L.
3       DuBois, Raymond N.
4 <120> TITLE OF INVENTION: Mammalian Genes Involved in Viral
5       Infection and Tumor Suppression
6 <130> FILE REFERENCE: 01123.0004
7 <140> CURRENT APPLICATION NUMBER: US/09/509,712A
8 <141> CURRENT FILING DATE: 2001-08-29
9 <150> PRIOR APPLICATION NUMBER: PCT/US98/21276
10 <151> PRIOR FILING DATE: 1998-10-08
11 <150> PRIOR APPLICATION NUMBER: 60/062,021
12 <151> PRIOR FILING DATE: 1997-10-10
13 <160> NUMBER OF SEQ ID NOS: 127
14 <170> SOFTWARE: FastSEQ for Windows Version 4.0
16 <210> SEQ ID NO: 1
17 <211> LENGTH: 925
18 <212> TYPE: DNA
19 <213> ORGANISM: Rattus norvegicus
20 <221> NAME/KEY: misc_feature
21 <222> LOCATION: 1- 925
22 <223> OTHER INFORMATION: n = g, a, c or t(u)
23 <400> SEQUENCE: 1
W--> 24  gggggaaaac cnggnaattg ttttttgacg anccaaaaag gggncnagna gcnnttntcc  60
W--> 25  tanatggggn cgggatcntn tccnaggana gattnatgga gtatnccttt tttgcncnaa 120
W--> 26  ggttgattgc tcttgaaagg ntttgaggtg naattcctcc gtnagtttga ccgtagtcgg 180
W--> 27  atntgaagag ggattgttna gcagncataa tttcattccc tgnacaccca gtaacnnttt 240
W--> 28  accgtcattt ggttgggaat tgatntcggg aggtancaan ggccacagtt atttattgtt 300
W--> 29  ncggaggatt gcaccaattn ggccggctgc ctctganatc tgtttctcat ccatgccggt 360
W--> 30  tcacccagac gaaagccgaa agcntcggga gtcctaactn tagtccntga aagtcattcc 420
     31  cagctgcgta attgggctgt gcagagtccc agctcggtaa atatttgccc cgtgactgag 480
     32  ctggagagaa tgctcctttc ttggtcctgg gcagctcttg gcagctcaca tgcactgttt 540
     33  acctatcctc ccacattccc ccctgaggaa tcatcgtgcc tcggttccct taagtcctct 600
     34  caacagaaaa caaggcagag tggaacgaag gaaagtgcgt ggccgttaga aagcctgtct 660
     35  cgaatctgtc ccacgtgcct caggtagcgt tccaaacagc aaagattcta gtgaagaaaa 720
     36  ataccgtccg gtcaattagt caggtggaca gagcaggacc cggtgtcttg gaagcctcgt 780
     37  ccattcctct ggggaaggtg gggggggcg tgtaatgcag ctctcaagaa gaaggtattt 840
     38  ttgttttcct ggagaaactg ccatcccagg agctgagagt ggatcagtag gaaggcctgt 900
     39  gacaggaagc agggaggttc agcng                                      925
41 <210> SEQ ID NO: 2
42 <211> LENGTH: 554
43 <212> TYPE: DNA
44 <213> ORGANISM: Rattus norvegicus
45 <221> NAME/KEY: misc_feature
46 <222> LOCATION: 1- 554
47 <223> OTHER INFORMATION: n = g, a, c or t(u)
48 <400> SEQUENCE: 2
49  caagatngan ggggcggcgg ttcgnccaga gagcgggtag ggaagggaac gcgccggatg  60
```

Does Not Comply
Corrected Diskette Needed

(global error)
insert this mandatory numeric identifier wherever <221>, <222>, or <223> is shown

<2207

<2207

RAW SEQUENCE LISTING DATE: 11/07/2001
PATENT APPLICATION: US/09/509,712A TIME: 13:41:54

Input Set : N:\Crf3\09192001\I509712A.raw
Output Set: N:\CRF3\11072001\I509712A.raw

```
W--> 50        agccngggtg cgganagcca gaccccaggc gtgggaaggg gagagagata gagcggccgg     120
W--> 51        ttgggaagag gaggaccgtg gttnataaat aacagaaagc ccagagggac gtanccatcc     180
W--> 52        gggatggaga gaggtaggga atccagntgt aagtcccaaa ctgccaccac cttcatnaga     240
W--> 53        actgcttcgt gtaaggtcac gcaccgggcc agctgtccng agtggcggtc ctggcgtgtt     300
W--> 54        aagttagcta aagtnactgc aactccgnct gtgcagactg ntcgtaaatt ctctctgtcc     360
W--> 55        gccaaattct ccctcctatt aaacttttca cttcctttca cttagtttcc tnacttcttt     420
W--> 56        caaacggaag ctgtaactga gcctgccacc cnganacntt gtggttgcca tttttatgct     480
W--> 57        aaagtaatcg tgtttttttat gcctgtcaac tccctttca tntaaagcag ggcntaccct     540
     58        attataactc tgcc                                                      554
     60 <210> SEQ ID NO: 3
     61 <211> LENGTH: 891
     62 <212> TYPE: DNA
     63 <213> ORGANISM: Rattus norvegicus                          >C2207
     64 <221> NAME/KEY: misc_feature
     65 <222> LOCATION: 1- ~~825~~ 878 (See below)
     66 <223> OTHER INFORMATION: n = g, a, c or t(u)
     67 <400> SEQUENCE: 3
W--> 68         ttngaaanaa tttccgtnaa ggtcngnaat nggccccgga aaaaatgngt tcctccccac      60
W--> 69         cttcattggn gcggatcctg ccngggaggc caatggttta acaaataatc tttnggagnt     120
W--> 70         ntggtngggg ggggagggac ncccacagan tcatgnggtg gttngggngg ngggcatcgt     180
W--> 71         tnngatatta tcacattntg ngaanctatg tnggggcttc ctttcngaca ggtggtggtt     240
W--> 72         nnacangngg atgtgtgctt ctttttcag cagtggtgga cccggattct aagacccttа      300
W--> 73         cngtaacaat gccctntttt cctaagccta accagtcctt tangaggant gctcttgggn     360
W--> 74         acccatgctg nntcacctag ccttggntca catnttnnac acaggaaaag gcagcatgtc     420
W--> 75         ttntnggagc tcagcttatt cccttcccnt cccatccagn atctccctgg gntggatgag     480
W--> 76         gtggatgacg catcttcaaa gcaccccacg tntcatggga tgtgcacagg agcttcgttg     540
W--> 77         gaaatgtgtt gcgcgaccag gcttgtgtag gaaacaacag actactcgaa attaaagtcn     600
W--> 78         taccttgcag ggttctcaga ggcttttacg cattaataaa catttgaatc ntaagaaggg     660
W--> 79         agcacagcat gtaatattnt tcaaattatc aggcnttgca accttcatta gtttctctta     720
W--> 80         cgcagctggg ngtggtggtg tgtaccttta atctcagcac tgaggaggca cngatatctc     780
W--> 81         catctctgtg acttccagac cggcntcgcc agagcaagtt ccaggccacc cagatgagat     840
W--> 82         gctcacagag gggacctttt tntgatgacc aacgnagnat gcaagtaagg a               891
     84 <210> SEQ ID NO: 4
     85 <211> LENGTH: 974
     86 <212> TYPE: DNA
     87 <213> ORGANISM: Rattus norvegicus                          >C2207
     88 <221> NAME/KEY: misc_feature
     89 <222> LOCATION: 1- 974
     90 <223> OTHER INFORMATION: n = g, a, c or t(u)
     91 <400> SEQUENCE: 4
W--> 92         aaaanaanat attccgnntc tnntagcnna gaagttntnc gagcnntccc ccgtntttttt      60
W--> 93         aaaaacccnc ggattccggn nntcgggntt taanngnttt tttaanggcc cnaagncccn     120
W--> 94         nttattgccg ncntttcccc cccgctnttg cnccccttta cttngagant ngtgntncna     180
W--> 95         agatttnaag gttnttgccc ccccggcttt tnttcccctn nttttccccn nagntttaaa     240
W--> 96         accggtntgg gttncnantt nnttgnancc nccnattggg gtttccgntt accngggttt     300
W--> 97         ttccccatgn ccgttccctc caatnttgna cttcccnggt cngggtccna atnccnngna     360
W--> 98         acngntcnan ccttattgac aattaatttt tccttgngna ntctgncccc cngnantttg     420
W--> 99         gggttcttgg gngcagggcc ttttttttcnt tggnngcaan cncataaatn ttaccagntt     480
W--> 100         gattgctaag gaagtancca tggttgngaa ccccccccttn ttntctccca gatggaaccc     540
```

6 last "n" is at location 878

RAW SEQUENCE LISTING DATE: 11/07/2001
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Input Set : N:\Crf3\09192001\I509712A.raw
Output Set: N:\CRF3\11072001\I509712A.raw

```
W--> 101        aggattttgg aactgcagag gcttcagggt cttgggaagc ggaggcagnn aaagattgga     600
W--> 102        gtgcactgtc cttttgcaat atggggtttg cctgcctgct ggctcntctc ctgctntntc     660
W--> 103        agatggtgac tgaggctact tcngcaggac tnggaataat catgtccagg tggctgccct     720
W--> 104        tccgagcaga aagggacaga cgtggggcga tgaagttgct atcgtttntt ttttttctg      780
     105        cacagactgc aaagtgtgca gagggaggga ggctgtgcaa aaaaaaaaaa aaaaaaaaaa     840
     106        aaaaaaaaaa ccgaggacgc agaagttaga ctgctgaccc atttggtgca tgtgtgccca     900
W--> 107        tggagggagg ggaccttntt taaagggttc acgcggcacg cantgggnaa nngnncctnt     960
W--> 108        acgnnnctcc caga                                                      974
     110 <210> SEQ ID NO: 5
     111 <211> LENGTH: 850
     112 <212> TYPE: DNA
     113 <213> ORGANISM: Rattus norvegicus
     114 <221> NAME/KEY: misc_feature
     115 <222> LOCATION: 1- 850
     116 <223> OTHER INFORMATION: n = g, a, c or t(u)
     117 <400> SEQUENCE: 5
W--> 118        antttccct caagnaaant ntggtttggg caacttgaag acgcttnnac cnaaaaccct      60
W--> 119        tgnggagntt ggngaccttn ttaccgnaan gagtgggaaa cgttttcctc cgggttnang     120
W--> 120        gttaggggga cccgnnggaa aattttaaaa ccnngngggc tttttcgaat taaggggaaa     180
W--> 121        ngcggtttng gtnnntgaag ggcgggnggt tggagtcnna gtccagagtt gatttccacc     240
W--> 122        cacaaatntg ggaggtgncg gggaatgntg ncnttttctt gngatgaggg ntgccgtncc     300
W--> 123        ggantaacag ngnttgcntt gtntngcnaa acgaagagtn tcctgnttgg aataggngtt     360
W--> 124        cngttcgang ganccagatt tangngntgg agnaaggatt nggcagataa angcntgaga     420
W--> 125        natgnancnt ggancaggtc nggncnnagn ntacagatga tgnncccana canganataa     480
W--> 126        ntncagatca cagtcgtacc cgnggctggg ccatgaanag ggcatcccca gacnnacaca     540
W--> 127        ngccttnana antgntcaga gaaccancag tggntanggg ntgcccnnnn naccagggaa     600
W--> 128        gacccggggc gtgncggata ttgacacanc agatnncatt tggggncggt tcgagggttn     660
W--> 129        atgntcnccg agtacnagan angatcntcc aacccggaat ncggtgctcc ngtcgtccga     720
W--> 130        tgnaatgagt cgnccggnaa cctcatatcc aagaaacnat acagcagtgg nntccgagtc     780
W--> 131        tcgtatantc nttgcgggng gaggctatnt tcagaggnca agattaccgt tagcgggana     840
W--> 132        aagtngaana                                                           850
     134 <210> SEQ ID NO: 6
     135 <211> LENGTH: 531
     136 <212> TYPE: DNA
     137 <213> ORGANISM: Rattus norvegicus
     138 <221> NAME/KEY: misc_feature
     139 <222> LOCATION: 1- 531
     140 <223> OTHER INFORMATION: n = g, a, c or t(u)
     141 <400> SEQUENCE: 6
W--> 142        ttgnggcngg gtctcctctg ngtgngngtn tccccnanag ggggggtctc acagtgtnng      60
W--> 143        ngtctnntgt ctgtgtngtg cccctgtccn catctctcac nccagggaga gagatgtgag     120
W--> 144        ananacatca gagatctctn gnacagtgtt tcacaagagt ctatcncana gagcacatct     180
W--> 145        gcccggggng anacacaact ctaaatgtgt ctcanntgat ctctctnttg tgtctctnac     240
W--> 146        atatgnggac atgctctcag agtatnggnt ctcttgngcn cttntgcaca cacacacaca     300
W--> 147        cacacacaca cacacacaca cacncttctc tctggcacag ggntatggca nagcacatnt     360
W--> 148        tnngagntca nagctntata tgagtgtgtg gcgaaaggng tnatnanann gacnncccca     420
W--> 149        gcnnatatag gggggngnnc tctngggctc tcttnggnaa tntgngggng agtctgcnca     480
W--> 150        cacaggcgct cnnacccanc nnnttggggc cccccaggng tttttcnccc c              531
     152 <210> SEQ ID NO: 7
```

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Input Set : N:\Crf3\09192001\I509712A.raw
Output Set: N:\CRF3\11072001\I509712A.raw

153 <211> LENGTH: 572
154 <212> TYPE: DNA
155 <213> ORGANISM: Rattus norvegicus >C2207
156 <221> NAME/KEY: misc_feature
157 <222> LOCATION: 1- 572
158 <223> OTHER INFORMATION: n = g, a, c or t(u)
159 <400> SEQUENCE: 7

W--> 160 ttttntgtg gcccttaaa ctctgngtgn ccgtntnccc nagaggggg gtctcacaag 60
W--> 161 gagacancgg nnacacagag gttttgngnn tattgngagt ctctgcgcac nccananttt 120
W--> 162 aaccncgggg nctcntgttt tattttaaaa aaaaagagtc ncatgtntat ttctctnatg 180
W--> 163 tgaaaatcnc attcanagtt ntggggtttc ccntgaggag anatagagtt tcacactctt 240
W--> 164 ctctccgagg ggtcntcnca tgtntctccc caatgtgngn ggnacacaca tgnggccccn 300
W--> 165 agggggtgng ctctctctgc ncagggcncc ccccaanang tagaganaca ntgtggtgtt 360
W--> 166 tcacaacaca attcncgaga nattntgttc cncantggnn gtctnagntc ncatgttgtg 420
W--> 167 gngacangtt agnncncccc atnttcnccc ccctttcaca ctgccccnag agagagaaan 480
W--> 168 tctnggcccc ctctanannt ntttttaaat cnccccnnac cacaggtntt cccagggtat 540
W--> 169 gngacntcnc cnnccccncn aaagatntgc nc 572

171 <210> SEQ ID NO: 8
172 <211> LENGTH: 906
173 <212> TYPE: DNA
174 <213> ORGANISM: Rattus norvegicus >C2207
175 <221> NAME/KEY: misc_feature
176 <222> LOCATION: 1- 906
177 <223> OTHER INFORMATION: n = g, a, c or t(u)
178 <400> SEQUENCE: 8

W--> 179 tgggagtctc tctcatatgg cgcnttcncc aaaggggngt ctctntccng agncgcanac 60
W--> 180 gcgagaanac tctgtnnant ngtctccccc cncnccnaca gngtganant caaaacctct 120
W--> 181 agagcccccc agaaancccc tntctcaaan aaagagaaag agaagancga gnagnagaga 180
W--> 182 gananagaga gagagagtgt gganctntnt cctcngancc ccannnanan ngtgnggcnc 240
W--> 183 actcncnngt gnngngnacc ccnggggatt tncgcgtgtc cccttgngct ctgtntanga 300
W--> 184 gananatatg tntagtctct ctntcgcccc ctccgntgtc acgtgtgcgg ggcccnngag 360
W--> 185 acacagacac ntctctcang gggaacacat anngactcnc acntgtgttt atattcnccc 420
W--> 186 ctcccnctca cacanacaca cacacagnag atattnngct actctctctc tgtcacaggg 480
W--> 187 gtacanattt antctnggcc anacccctct cngaagngng ggcanngtaa accccgcccc 540
W--> 188 ctctcngaga angngagggc gntttacntt cccngtggcg tgtncgngcc cccgagactc 600
W--> 189 cccttngnac ccccctntna accctctntt tgaacncaac ncaccntccc cnttttctcg 660
W--> 190 gggnnggncc ngcncccnct ctcncaaaaa aaattnnaan ttngtcccct nccccnttnt 720
W--> 191 ttcnggnana aaccgtgtcc gggggggggan nactcttttt tgnccttaaa atcaanttt 780
W--> 192 ttccccttt ccnggggacc cccgnnttcc tttttaaaaa aaaanaaccc tttctccctt 840
W--> 193 ttaaaagnac ccnttttttc naaaaccgtt ccgnatttaa ttcctaaatt cccttccccn 900
W--> 194 ncccgg 906

196 <210> SEQ ID NO: 9
197 <211> LENGTH: 914
198 <212> TYPE: DNA
199 <213> ORGANISM: Rattus norvegicus >C2207
200 <221> NAME/KEY: misc_feature
201 <222> LOCATION: 1- 914
202 <223> OTHER INFORMATION: n = g, a, c or t(u)
203 <400> SEQUENCE: 9

RAW SEQUENCE LISTING DATE: 11/07/2001
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Input Set : N:\Crf3\09192001\I509712A.raw
Output Set: N:\CRF3\11072001\I509712A.raw

```
W--> 204  gggatgngcc ctcagatcaa tacacccctc ngggggngtc tctctctatc tcccncagna   60
W--> 205  gactcccatc tctntntntn cccccaganc tggngaacgg ngtgtggnga nccntntctg  120
W--> 206  ttctcnantc tctaaaagng cnaaaagcgc ananacacgn gcctctctat anatctcacg  180
W--> 207  tgtcccnngn nctctcngac ccctnntctg tntgagagac accctntctc aaaatatagt  240
W--> 208  gtacacgngc tttgnggctc tccccttttc tctccactnt tgagngngaa acgcggngtt  300
W--> 209  ntctctgaga tgtaganagn gtcccctnct cnatatatgt gttncccact ccnnagggng  360
W--> 210  tctcataaaa atcncntntc tcaacaccac cncctcnacc ccccncacga gaacacntcn  420
W--> 211  ccaccncnan gacacaaana naaggngtnn anaaccccan aaaaactnng ntntcngntt  480
W--> 212  tacacacaca cacacncacn ctcncncaca cccccacnna aatgggagaa aaaacagaga  540
W--> 213  ggngtgggtg ttngnntcaa caccntntta cctctctgnt gnnanttgag aaaatatttc  600
W--> 214  tntncttacc cctctcccct ctctgtgtgt ngannatatc ngntctagat gtcctnaccc  660
W--> 215  tccccaaacc tttctcnggn agagacntct ctntntttt cccccncttc catttgaaan  720
W--> 216  anangagaag gnccaaaaag gngggngtct tctcgggaat ncncccttt ggcccccaa   780
W--> 217  cctgggtttt tttccccctt ccttttaatn anttttcna nacaaanctt tnngngtttn  840
W--> 218  ggaaaangcc tttnnctgnn nntttttcc cttcccctt tnnangggnt tcccccccc    900
W--> 219  ccngaatttt tttt                                                   914
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221 <210> SEQ ID NO: 10
222 <211> LENGTH: 400
223 <212> TYPE: DNA
224 <213> ORGANISM: Rattus norvegicus ——> (2207
225 <221> NAME/KEY: misc_feature
226 <222> LOCATION: 1- 400
227 <223> OTHER INFORMATION: n = g, a, c or t(u)
228 <400> SEQUENCE: 10

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229  ttcctgggtg cggtctcctc tgagatagtg tatcccctat agggggggtc tcactttagc   60
230  acagtttatg aatattatta catatttcac aagactttat attgttataa tatgcctcat  120
231  gtgagatata tgtgattctg tggtggtgtt ctcagagggg gtttgggtta ttggggataa  180
232  tagtttgccc ctcgcggggt ctatatttat atatgtgaca caatatatta gagagatttt  240
233  tggttatata tatttccctt cgcgggggtg gagatttatc acagggggag agcttttccc  300
234  ttgttagcaa aagtccctgg tctcgtcccc catctcccaa aaaaaaaaaa atgtgaaaaa  360
235  aaaaaaaaaa agggcccctc ttgagtgatg tccccttctt                       400
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237 <210> SEQ ID NO: 11
238 <211> LENGTH: 880
239 <212> TYPE: DNA
240 <213> ORGANISM: Rattus norvegicus ——> (2207
241 <221> NAME/KEY: misc_feature
242 <222> LOCATION: 1- 880
243 <223> OTHER INFORMATION: n = g, a, c or t(u)
244 <400> SEQUENCE: 11

```
W--> 245  acccaatctt nanggtggca gtgnggnnga tcttaacggt ttttnagaaa aaaaantnct   60
W--> 246  tcgctcncac ccccaagcct cccnttctta ncagcttttt tatangaaaa aagatgataa  120
W--> 247  cgaaatttta aaaaccgtcg ttagaggaaa tgaaggttca gccgaccatt acctganagt  180
W--> 248  aatgaaggtn ttccggaggg ttgccttcca atcccagatg gatttgagtt tcaggatcaa  240
W--> 249  ttcagttacc gntgaccatc caccnncctc cngtataatc attngatgag gatgaatggt  300
W--> 250  gagtgagtga tgatgatgat gatgatgatg aagggatgag aagnacacta tgataacaag  360
W--> 251  tgtctcagtc cacattaagg tttgcctgna aattagtgca taagccatgg gagacaaatt  420
W--> 252  cttttcnnac acaattaata gtntcttant ccttcccatc ttctctgccc cattctgttt  480
      253  tccaccacag gtctgcagcg ggctacagct tccagtctcc aagcaaatac cagaactgga  540
      254  ggagaaaatt ccagtccagt gagtcatggg caggggggagg ggtggggtaa gggcagtggc  600
```

The types of errors shown exist throughout the Sequence Listing. Please check subsequent sequences for similar errors.

FYI Use of n and/or Xaa has been detected in the Sequence Listing. Review the Sequence Listing to insure a corresponding explanation is presented in the <220> to <223> fields of each sequence using n or Xaa.

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Input Set : N:\Crf3\09192001\I509712A.raw
Output Set: N:\CRF3\11072001\I509712A.raw

L:7 M:270 C: Current Application Number differs, Wrong Format
L:24 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:24 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:25 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:25 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:26 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:26 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:27 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:27 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:28 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:28 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:29 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:29 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:30 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:30 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:39 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:39 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:49 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2
L:49 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:50 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2
L:50 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:51 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2
L:51 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:52 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2
L:52 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:53 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2
L:53 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:54 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2
L:54 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:55 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2
L:55 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:56 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2
L:56 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:57 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2
L:57 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:68 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:3
L:68 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:69 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:3
L:69 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:70 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:3
L:70 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:71 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:3
L:71 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:72 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:3
L:72 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:73 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:3
L:73 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:74 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:3

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Input Set : N:\Crf3\09192001\I509712A.raw
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L:74 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:75 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:3
L:75 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:76 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:3
L:76 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:77 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:3
L:77 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:78 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:3
L:78 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:79 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:3
L:79 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:80 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:3
L:80 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:81 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:3
L:81 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:82 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:3
L:82 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:92 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:4
L:92 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:93 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:4
L:93 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:94 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:4
L:94 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:95 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:4
L:95 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:96 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:4
L:96 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:97 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:4
L:97 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:98 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:4
L:98 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:99 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:4
L:99 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:100 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:4
L:100 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:101 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:4
L:101 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:102 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:4
L:102 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:103 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:4
L:103 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:104 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:4
L:104 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:107 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:4
L:107 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:108 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:4
L:108 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:118 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:5
L:118 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5

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Input Set : N:\Crf3\09192001\I509712A.raw
Output Set: N:\CRF3\11072001\I509712A.raw

L:119 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:5
L:119 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:120 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:5
L:120 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5